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FIG. 1A

TRQSP-----RLPFTSPSL--SLCLSSNRVHLLAAASKRDQMAPLGDG-GAAAAAASN  
-ARASPPHPKK--FPTIPSLSLNLDLSLSPVLFARSR-----VVVGGGAMAGRS  
HERTSLSCRSKTVFQNPNTLSLP-----MALNNGVPATGT  
-----MAPLGDAVAPAAAAAP  
-----MAVNG-----ASSS  
1 60

\* \* \* \* \*  
NLVVSFGEMLIDFVPDVAGLSLAESGGFVKAPGGAPANVACAIKLGSSAFVKGFGDDE  
ELVVSFGEMLIDFVPTVAGVSLAEAPAFVVKAPGGAPANVAIAVARLGGAAAFVKGKLGDDDE  
GLVVSFGEMLIDFVPTVSGVSLAEAPGFLKAPGGAPANVAIAVSRLLGGKAAAFVKGKLGDDDE  
GLVVSFGEMLIDFVPDVAGVSLAEESGGFVKAPGGAPANVACAIKLGSSAFVKGFGDDE  
GLVVSFGEMLIDFVPTVSGVSLAEAPGFLKAPGGAPANVAIAVTRLLGGKSAFVKGKLGDDDE  
61 120

\* \* \* \* \*  
FGHMLVNLKQNNVNSEGCLFDKHARTALAFVTLKHDGEREFMFYRNPSADMLLTHEAELD  
FGRMLAAILRDNGVDDGGVFDAGARTALAFVTLRADGEREFMFYRNPSADMLLTHAELN  
FGHMLAGILKENGVRADGINFDQGARTALAFVTLRADGEREFMFYRNPSADMLLKPEELN  
FGHMLVEILKQNGVNAEGCLFDQHARTALAFVTLKSDGEREFMFYRNPSADMLLTHEAELN  
FGHMLAGILKTNGVQAEGINFDKGARTALAFVTLRADGEREFMFYRNPSADMLLTPAELN  
121 180

\* \* \* \* \*  
LGLVRRRAKVHYGSIISLISEPCRSAAHMAAMRAAKAAGVLCSDPNVRLPLWSPDAAREG  
VELIKRAAVFHYGSIISLIAEPCRSAAHLRAAMEIAKEAGALLSYDPNLRALWPSREEARTK  
LELIRSAKVHYGSIISLIVEPCRSAAHLKAMEVAKESGCLLSYDPNLRPLWPSAEERKQ  
LDLIRRAIRFHYGSIISLITEPCRSAAHVAATRAAKSAGILCSYDPNVRPLWPSAQAAARDG  
LDLIRSAKVHYGSIISLIVEPCRAAHKAMEVAKEAGALLSYDPNLRPLWPSAEERKQ  
181 240

FIG. 1A

[illegible]

SEQ	ID	NO:02
SEQ	ID	NO:06
SEQ	ID	NO:08
SEQ	ID	NO:12
SEQ	ID	NO:13

SEQ	ID NO:02
SEQ	ID NO:06
SEQ	ID NO:08
SEQ	ID NO:12
SEQ	ID NO:13

SEQ	ID	NO:02
SEQ	ID	NO:06
SEQ	ID	NO:08
SEQ	ID	NO:12
SEQ	ID	NO:13

**FIG. 1B**